SEQUENCE LISTING

<110> Turner, C. Alexander Jr.
Donoho, Gregory
Nehls, Michael
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<120> Novel Human Proteins and Polynucleotides
Encoding the Same

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<150> US 60/160,106
<151> 1999-10-18

<150> US 60/162,547
<151> 1999-10-29

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Cys Leu Phe His Gly Lys Arg Tyr Ser Pro Gly Glu Ser Trp His Pro
        35                  40                  45
Tyr Leu Glu Pro Gln Gly Leu Met Tyr Cys Leu Arg Cys Thr Cys Ser
    50                  55                  60
Glu Gly Ala His Val Ser Cys Tyr Arg Leu His Cys Pro Pro Val His
65                  70                  75                  80
Cys Pro Gln Pro Val Thr Glu Pro Gln Gln Cys Cys Pro Lys Cys Val
                85                  90                  95
Glu Pro His Thr Pro Ser Gly Leu Arg Ala Pro Pro Lys Ser Cys Gln
            100                 105                 110
His Asn Gly Thr Met Tyr Gln His Gly Glu Ile Phe Ser Ala His Glu
        115                 120                 125
Leu Phe Pro Ser Arg Leu Pro Asn Gln Cys Val Leu Cys Ser Cys Thr
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Glu Gly Gln Ile Tyr Cys Gly Leu Thr Thr Cys Pro Glu Pro Gly Cys
145                 150                 155                 160
Pro Ala Pro Leu Pro Leu Pro Asp Ser Cys Cys Gln Ala Cys Lys Asp
                165                 170                 175
Glu Ala Ser Glu Gln Ser Asp Glu Glu Asp Ser Val Gln Ser Leu His
            180                 185                 190
Gly Val Arg His Pro Gln Asp Pro Cys Ser Ser Asp Ala Gly Arg Lys
        195                 200                 205
Arg Gly Pro Gly Thr Pro Ala Pro Thr Gly Leu Ser Ala Pro Leu Ser
    210                 215                 220
Phe Ile Pro Arg His Phe Arg Pro Lys Gly Ala Gly Ser Thr Thr Val
225                 230                 235                 240
Lys Ile Val Leu Lys Glu Lys His Lys Lys Ala Cys Val His Gly Gly
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Lys Thr Tyr Ser His Gly Glu Val Trp His Pro Ala Phe Arg Ala Phe
            260                 265                 270
Gly Pro Leu Pro Cys Ile Leu Cys Thr Cys Glu Asp Gly Arg Gln Asp
        275                 280                 285
Cys Gln Arg Val Thr Cys Pro Thr Glu Tyr Pro Cys Arg His Pro Glu
    290                 295                 300
Lys Val Ala Gly Lys Cys Cys Lys Ile Cys Pro Glu Asp Lys Ala Asp
305                 310                 315                 320
Pro Gly His Ser Glu Ile Ser Ser Thr Arg Cys Pro Lys Ala Pro Gly
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Arg Val Leu Val His Thr Ser Val Ser Pro Ser Pro Asp Asn Leu Arg
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Arg Phe Ala Leu Glu His Glu Ala Ser Asp Leu Val Glu Ile Tyr Leu
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Trp Lys Leu Val Lys Asp Glu Glu Thr Glu Ala Gln Arg Gly Glu Val
    370                 375                 380
Pro Gly Pro Arg Pro His Ser Gln Asn Leu Pro Leu Asp Ser Asp Gln
385                 390                 395                 400
Glu Ser Gln Glu Ala Arg Leu Pro Glu Arg Gly Thr Ala Leu Pro Thr
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                405                 410                 415
Ala Arg Trp Pro Pro Arg Arg Ser Leu Glu Arg Leu Pro Ser Pro Asp
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<213> homo sapiens

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gcccctctga gcttcatccc tcgccacttc agacccaagg gagcaggcag cacaactgtc     720
aagatcgtcc tgaaggagaa acataagaaa gcctgtgtgc atggcgggaa gacgtactcc     780
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cgtcaccccg agaaagtggc tgggaagtgc tgcaagattt gcccagagga caaagcagac     960
cctggccaca gtgagatcag ttctaccagg tgtcccaagg caccgggccg ggtcctcgtc    1020
cacacatcgg tatccccaag cccagacaac ctgcgtcgct ttgccctgga acacgaggcc    1080
tcggacttgg tggagatcta cctctggaag ctggtraaag gaatcttcca cttgactcag    1140
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<210> 4
<211> 429
<212> PRT
<213> homo sapiens

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            20                  25                  30
Cys Leu Phe His Gly Lys Arg Tyr Ser Pro Gly Glu Ser Trp His Pro
        35                  40                  45
Tyr Leu Glu Pro Gln Gly Leu Met Tyr Cys Leu Arg Cys Thr Cys Ser
    50                  55                  60
Glu Gly Ala His Val Ser Cys Tyr Arg Leu His Cys Pro Pro Val His
65                  70                  75                  80
Cys Pro Gln Pro Val Thr Glu Pro Gln Gln Cys Cys Pro Lys Cys Val
                85                  90                  95
Glu Pro His Thr Pro Ser Gly Leu Arg Ala Pro Pro Lys Ser Cys Gln
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            100                 105                 110
His Asn Gly Thr Met Tyr Gln His Gly Glu Ile Phe Ser Ala His Glu
        115                 120                 125
Leu Phe Pro Ser Arg Leu Pro Asn Gln Cys Val Leu Cys Ser Cys Thr
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Glu Gly Gln Ile Tyr Cys Gly Leu Thr Thr Cys Pro Glu Pro Gly Cys
145                 150                 155                 160
Pro Ala Pro Leu Pro Leu Pro Asp Ser Cys Cys Gln Ala Cys Lys Asp
                165                 170                 175
Glu Ala Ser Glu Gln Ser Asp Glu Glu Asp Ser Val Gln Ser Leu His
            180                 185                 190
Gly Val Arg His Pro Gln Asp Pro Cys Ser Ser Asp Ala Gly Arg Lys
        195                 200                 205
Arg Gly Pro Gly Thr Pro Ala Pro Thr Gly Leu Ser Ala Pro Leu Ser
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Phe Ile Pro Arg His Phe Arg Pro Lys Gly Ala Gly Ser Thr Thr Val
225                 230                 235                 240
Lys Ile Val Leu Lys Glu Lys His Lys Lys Ala Cys Val His Gly Gly
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Lys Thr Tyr Ser His Gly Glu Val Trp His Pro Ala Phe Arg Ala Phe
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Gly Pro Leu Pro Cys Ile Leu Cys Thr Cys Glu Asp Gly Arg Gln Asp
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Cys Gln Arg Val Thr Cys Pro Thr Glu Tyr Pro Cys Arg His Pro Glu
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Lys Val Ala Gly Lys Cys Cys Lys Ile Cys Pro Glu Asp Lys Ala Asp
305                 310                 315                 320
Pro Gly His Ser Glu Ile Ser Ser Thr Arg Cys Pro Lys Ala Pro Gly
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Arg Val Leu Val His Thr Ser Val Ser Pro Ser Pro Asp Asn Leu Arg
            340                 345                 350
Arg Phe Ala Leu Glu His Glu Ala Ser Asp Leu Val Glu Ile Tyr Leu
        355                 360                 365
Trp Lys Leu Val Lys Gly Ile Phe His Leu Thr Gln Ile Lys Lys Val
    370                 375                 380
Arg Lys Gln Asp Phe Gln Lys Glu Ala Gln His Phe Arg Leu Leu Ala
385                 390                 395                 400
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<212> PRT
<213> Homo sapiens

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His Thr Tyr Pro Arg Asn Thr Val Leu Val Trp Arg Leu Val Ala Val
65                  70                  75                  80
Glu Glu Asn Val Trp Ile Gln Leu Thr Phe Asp Glu Arg Phe Gly Leu
                85                  90                  95
Glu Asp Pro Glu Asp Asp Ile Cys Lys Tyr Asp Phe Val Glu Val Glu
            100                 105                 110
Glu Pro Ser Asp Gly Thr Ile Leu Gly Arg Trp Cys Gly Ser Gly Thr
        115                 120                 125
Val Pro Gly Lys Gln Ile Ser Lys Gly Asn Gln Ile Arg Ile Arg Phe
    130                 135                 140
Val Ser Asp Glu Tyr Phe Pro Ser Glu Pro Gly Phe Cys Ile His Tyr
145                 150                 155                 160
Asn Ile Val Met Pro Gln Phe Thr Glu Ala Val Ser Pro Ser Val Leu
                165                 170                 175
Pro Pro Ser Ala Leu Pro Leu Asp Leu Leu Asn Asn Ala Ile Thr Ala
            180                 185                 190
Phe Ser Thr Leu Glu Asp Leu Ile Arg Tyr Leu Glu Pro Glu Arg Trp
        195                 200                 205
Gln Leu Asp Leu Glu Asp Leu Tyr Arg Pro Thr Trp Gln Leu Leu Gly
    210                 215                 220
Lys Ala Phe Val Phe Gly Arg Lys Ser Arg Gly Asp Asn Tyr Glu Lys
225                 230                 235                 240
Glu Lys Asn Leu Lys Thr Asn Phe Tyr Lys Tyr Leu Ala Glu Gly Asn
                245                 250                 255
Phe Phe Asn Ile Ile Ile Val Lys Leu Phe Lys Lys Tyr Pro Leu Val
            260                 265                 270
His Tyr Leu Ser Phe Phe Phe Ser Phe Phe Ser Leu Phe Ala Phe Pro
        275                 280                 285
Pro Gln Lys Tyr Tyr Thr Met Phe Gln Glu Cys Met Thr Tyr Asp Leu
    290                 295                 300
Thr
305
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